

GenCore version 4.5  
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OM nucleic - nucleic search, using sw model

Run on: March 9, 2002, 00:54:06 ; Search time 319.49 Seconds  
(without alignments)  
17.722 Million cell updates/sec

Title: US-09-851-670-4

Perfect score: 25

Sequence: 1 acagtagcagcagcagcagcagc 25

Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 11323899 residues

Total number of hits satisfying chosen parameters: 515962

Minimum DB seq length: 0  
Maximum DB seq length: 60

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database :

1: ./cgn2\_6/ptodata/2/ina/5A.COMB.seq:\*  
2: ./cgn2\_6/ptodata/2/ina/6A.COMB.seq:\*  
3: ./cgn2\_6/ptodata/2/ina/6A.COMB.seq:\*  
4: ./cgn2\_6/ptodata/2/ina/6B.COMB.seq:\*  
5: ./cgn2\_6/ptodata/2/ina/PCITUS.COMB.seq:\*  
6: ./cgn2\_6/ptodata/2/ina/Backfiles1.seq:\*

Pred. NO. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match Length	ID	Description
1	16.2	64.8	57 4 US-09-043-303-14	Sequence 14, Appl
2	15.8	63.2	52 4 US-09-332-769-3	Sequence 3, Appl
3	15.8	63.2	52 4 US-09-456-153-3	Sequence 3, Appl
4	15.6	62.4	43 2 US-08-531-927B-17	Sequence 17, Appl
5	15	60.0	40 2 US-08-628-422-35	Sequence 35, Appl
6	14.6	58.4	21 2 US-08-863-639A-28	Sequence 28, Appl
7	14.6	58.4	21 2 US-08-863-639A-60	Sequence 60, Appl
8	14.6	58.4	24 2 US-08-863-639A-94	Sequence 94, Appl
9	14.6	58.4	30 1 US-08-068-747-6	Sequence 6, Appl
10	14.6	58.4	30 1 US-08-068-747-11	Sequence 11, Appl
11	14.6	58.4	30 2 US-08-863-639A-30	Sequence 30, Appl
12	14.6	58.4	30 4 US-09-135-994-4	Sequence 4, Appl
13	14.6	58.4	31 2 US-08-570-155-14	Sequence 14, Appl
14	14.6	58.4	31 5 PCT-US95-02861-14	Sequence 14, Appl
15	14.6	58.4	33 2 US-08-863-639A-29	Sequence 29, Appl
16	14.6	58.4	36 2 US-08-863-639A-31	Sequence 31, Appl
17	14.6	58.4	40 2 US-08-411-607A-7	Sequence 7, Appl
18	14.6	58.4	51 1 US-08-068-747-1	Sequence 1, Appl
19	14.4	57.6	42 4 US-09-142-355B-11	Sequence 11, Appl
20	14.2	56.8	23 1 US-08-390-850-363	Sequence 363, App
21	14.2	56.8	23 1 US-08-435-634-363	Sequence 363, App
22	14.2	56.8	24 1 US-08-423-383-76	Sequence 76, Appl
23	14.2	56.8	24 1 US-08-620-467A-43	Sequence 43, Appl
24	14.2	56.8	24 1 US-08-348-572-44	Sequence 44, Appl
25	14.2	56.8	24 2 US-08-437-353A-76	Sequence 76, Appl
26	14.2	56.8	24 3 US-08-559-205-47	Sequence 47, Appl
27	14.2	56.8	24 3 US-09-041-090B-44	Sequence 44, Appl

c 28	14.2	56.8	31 1	US-08-390-850-93	Sequence 93, Appl
c 29	14.2	56.8	31 1	US-08-390-850-94	Sequence 94, Appl
c 30	14.2	56.8	31 1	US-08-390-850-95	Sequence 95, Appl
c 31	14.2	56.8	31 1	US-08-435-634-93	Sequence 93, Appl
c 32	14.2	56.8	31 1	US-08-435-634-94	Sequence 94, Appl
c 33	14.2	56.8	31 1	US-08-435-634-95	Sequence 95, Appl
c 34	14	56.0	39 1	US-08-120-607A-5	Sequence 5, Appl
c 35	14	56.0	39 2	US-08-453-848-5	Sequence 5, Appl
c 36	14	56.0	39 4	US-09-169-027-5	Sequence 5, Appl
c 37	13.8	55.2	18 2	US-08-863-639A-17	Sequence 17, Appl
c 38	13.8	55.2	20 3	US-09-418-641-21	Sequence 21, Appl
c 39	13.8	55.2	20 3	US-09-418-641-80	Sequence 80, Appl
c 40	13.8	55.2	21 2	US-08-267-803B-66	Sequence 66, Appl
c 41	13.8	55.2	21 2	US-08-863-639A-40	Sequence 40, Appl
c 42	13.8	55.2	21 2	US-08-863-639A-66	Sequence 66, Appl
c 43	13.8	55.2	21 2	US-08-863-639A-69	Sequence 69, Appl
c 44	13.8	55.2	21 2	US-08-863-639A-87	Sequence 87, Appl
c 45	13.8	55.2	21 3	US-08-781-891-8	Sequence 8, Appl

#### ALIGNMENTS

```
RESULT 1
US-09-043-303-14
; Sequence 14, Application US/09043303
; Patent No. 6251389
; GENERAL INFORMATION:
; APPLICANT: SANPEI, Kazuhiro
; APPLICANT: TSUJI, Shoji
; TITLE OF INVENTION: Method for Diagnosing Spino cerebellar Ataxia Type 2 and
; TITLE OF INVENTION: Primers Therefor
; FILE REFERENCE: 0760-0241P
; CURRENT APPLICATION NUMBER: US/09/043,303
; CURRENT FILING DATE: 1998-05-18
; EARLIER APPLICATION NUMBER: PCT/JP96/01999
; EARLIER FILING DATE: 1996-07-18
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
; LENGTH: 57
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-043-303-14

Query Match      64.8%; Score 16.2; DB 4; Length 57;
Best Local Similarity 85.7%; Pred. No. 1.5e+02;
Matches 18; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY      2 cagtagcagcagcagcagcag 22
Db      10 cagcagcagcagcagcagcag 30

RESULT 2
US-09-332-769-3/c
; Sequence 3, Application US/09332769
; Patent No. 6172076
; GENERAL INFORMATION:
; APPLICANT: Embrey, Mark W.
; APPLICANT: Perlow, Debra S.
; APPLICANT: Wal, John S.
; APPLICANT: Hoffman, Jacob M.
; TITLE OF INVENTION: INHIBITORS OF PRENYL-PROTEIN
; TITLE OF INVENTION: TRANSFERASE
; FILE REFERENCE: 19982Y
; CURRENT APPLICATION NUMBER: US/09/332,769
; CURRENT FILING DATE: 1999-06-14
; EARLIER APPLICATION NUMBER: US 60/089,311
; EARLIER FILING DATE: 1998-06-15
; NUMBER OF SEQ ID NOS: 21
; SOFTWARE: FastSeq for Windows Version 3.0
```

SEQ ID NO 3  
LENGTH: 52  
TYPE: DNA  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: completely synthesized  
US-09-332-769-3

Query Match 63.2%; Score 15.8; DB 4; Length 52;  
Best Local Similarity 89.5%; Pred. No. 2.2e+02;  
Matches 17; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatg 20  
||| ||||| ||||| |||||  
DB 43 CAGCAGCAGCAGCAGCATG 25

RESULT 3  
US-09-456-153-3/C  
Sequence 3, Application US/09456153  
Patent No. 6284755  
GENERAL INFORMATION:  
APPLICANT: desolms, S. Jane  
APPLICANT: Graham, Samuel L.  
APPLICANT: Shaw, Anthony W.  
APPLICANT: Ciccotone, Terrence M.  
APPLICANT: Stokker, Gerald E.  
TITLE OF INVENTION: INHIBITORS OF PRENYL-PROTEIN  
FILE REFERENCE: 20312Y  
CURRENT APPLICATION NUMBER: US/09/456,153  
EARLIER FILING DATE: 1999-12-07  
EARLIER APPLICATION NUMBER: US 60/111,416  
EARLIER FILING DATE: 1998-12-08  
EARLIER APPLICATION NUMBER: US 60/129,282  
EARLIER FILING DATE: 1999-04-14  
NUMBER OF SEQ ID NOS: 21  
SOFTWARE: FastSeq. for Windows Version 3.0  
SEQ ID NO 3  
LENGTH: 52  
TYPE: DNA  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: completely synthesized  
US-09-456-153-3

Query Match 63.2%; Score 15.8; DB 4; Length 52;  
Best Local Similarity 89.5%; Pred. No. 2.2e+02;  
Matches 17; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatg 20  
||| ||||| ||||| |||||  
DB 43 CAGCAGCAGCAGCAGCATG 25

RESULT 4  
US-08-531-927B-17/C  
Sequence 17, Application US/08531927B  
Patent No. 5840491  
GENERAL INFORMATION:  
APPLICANT: Kakizuka, Akira  
TITLE OF INVENTION: DNA Sequence Encoding the Machado-Joseph  
Patent No. 5840491  
NUMBER OF SEQUENCES: 23  
CORRESPONDENCE ADDRESS:  
ADDRESS: Hamilton, Brook, Smith & Reynolds, P.C.  
STREET: Two Millitia Drive  
CITY: Lexington  
STATE: Massachusetts  
COUNTRY: USA

ZIP: 02173-4799  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/531,927B  
FILING DATE: 21-SEP-1995  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: JP H6-251600  
FILING DATE: 21-SEP-1994  
ATTORNEY/AGENT INFORMATION:  
NAME: Granahan, Patricia  
REGISTRATION NUMBER: 32,227  
REFERENCE/DOCKET NUMBER: A795-01A  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 617-861-9540  
TELEFAX: 617-861-6240  
INFORMATION FOR SEQ ID NO: 17:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 43 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: DNA (genomic)  
US-08-531-927B-17

Query Match 62.4%; Score 15.6; DB 2; Length 43;  
Best Local Similarity 81.8%; Pred. No. 2.6e+02;  
Matches 18; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatg 23  
||| ||||| ||||| |||||  
DB 26 CAGCAGCAGCAGCAGCAGA 5

RESULT 5  
US-08-628-422-35  
Sequence 35, Application US/08628422  
Patent No. 5837854  
GENERAL INFORMATION:  
APPLICANT: Mulder, Carel  
TITLE OF INVENTION: OLIGONUCLEOTIDES WITH ANTI-EPSTEIN-BARR  
NUMBER OF SEQUENCES: 63  
CORRESPONDENCE ADDRESS:  
ADDRESS: Fish & Richardson P.C.  
STREET: 225 Franklin Street  
CITY: Boston  
STATE: MA  
COUNTRY: USA  
ZIP: 02110-2804  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/628,422  
FILING DATE:  
CLASSIFICATION: 514  
ATTORNEY/AGENT INFORMATION:  
NAME: Fasse, J. Peter  
REGISTRATION NUMBER: 32,983  
REFERENCE/DOCKET NUMBER: 04020/094001  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 617/542-5070  
TELEFAX: 617/542-8906  
TELEX: 200154  
INFORMATION FOR SEQ ID NO: 35:

SEQUENCE CHARACTERISTICS:  
 LENGTH: 40 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: DNA  
 US-08-628-422-35

Query Match 60.0%; Score 15; DB 2; Length 40;  
 Best Local Similarity 78.3%; Pred. No. 4.5e+02;  
 Matches 18; Conservative 0; Mismatches 5; Indels 0; Gaps 0;

OY 1 acagtagcagcagacagcatgag 23  
 ||||| ||||| ||||| |||||  
 Db 3 ACAGTAGCGCCAGAGAGAGAGAG 25

RESULT 6  
 US-08-863-639A-28  
 Sequence 28, Application US/08863639A  
 Patent No. 5981185

GENERAL INFORMATION:  
 APPLICANT: Matson, Robert S.  
 APPLICANT: Coassin, Peter J.  
 APPLICANT: Rampal, Jang B.  
 APPLICANT: Caskey, C. T.  
 TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS  
 NUMBER OF SEQUENCES: 95  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: Sheldon & Mak  
 STREET: 225 South Lake Avenue, 9th Floor  
 CITY: Pasadena  
 STATE: CA  
 COUNTRY: USA  
 ZIP: 91101  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MB storage  
 COMPUTER: IBM compatible  
 OPERATING SYSTEM: Windows 95  
 SOFTWARE: Corel WordPerfect 8 version  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/863,639A  
 FILING DATE: May 28, 1997  
 CLASSIFICATION: 435  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Joseph E. Mueth  
 REGISTRATION NUMBER: 20,532  
 REFERENCE/DOCKET NUMBER: 11859-1  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (626) 796-4000  
 TELEFAX: (626) 795-6321  
 INFORMATION FOR SEQ ID NO: 28:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 21 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: Other nucleic acid  
 US-08-863-639A-28

Query Match 58.4%; Score 14.6; DB 2; Length 21;  
 Best Local Similarity 81.0%; Pred. No. 6.1e+02;  
 Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

OY 2 cagtagcagcagacagcatgag 22  
 ||| ||||| ||||| |||||  
 Db 1 CAGCAGCAGCAGCAGCAGCAG 21

RESULT 7  
 US-08-863-639A-60/c

Sequence 60, Application US/08863639A  
 Patent No. 5981185

GENERAL INFORMATION:  
 APPLICANT: Matson, Robert S.  
 APPLICANT: Coassin, Peter J.  
 APPLICANT: Rampal, Jang B.  
 APPLICANT: Caskey, C. T.  
 TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS  
 NUMBER OF SEQUENCES: 95  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: Sheldon & Mak  
 STREET: 225 South Lake Avenue, 9th Floor  
 CITY: Pasadena  
 STATE: CA  
 COUNTRY: USA  
 ZIP: 91101  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MB storage  
 COMPUTER: IBM compatible  
 OPERATING SYSTEM: Windows 95  
 SOFTWARE: Corel WordPerfect 8 version  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/863,639A  
 FILING DATE: May 28, 1997  
 CLASSIFICATION: 435  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Joseph E. Mueth  
 REGISTRATION NUMBER: 20,532  
 REFERENCE/DOCKET NUMBER: 11859-1  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (626) 796-4000  
 TELEFAX: (626) 795-6321  
 INFORMATION FOR SEQ ID NO: 60:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 21 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: Other nucleic acid  
 US-08-863-639A-60

Query Match 58.4%; Score 14.6; DB 2; Length 21;  
 Best Local Similarity 81.0%; Pred. No. 6.1e+02;  
 Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

OY 2 cagtagcagcagacagcatgag 22  
 ||| ||||| ||||| |||||  
 Db 21 CAGCAGCAGCAGCAGCAGCAG 1

RESULT 8  
 US-08-863-639A-94  
 Sequence 94, Application US/08863639A  
 Patent No. 5981185

GENERAL INFORMATION:  
 APPLICANT: Matson, Robert S.  
 APPLICANT: Coassin, Peter J.  
 APPLICANT: Rampal, Jang B.  
 APPLICANT: Caskey, C. T.  
 TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS  
 NUMBER OF SEQUENCES: 95  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: Sheldon & Mak  
 STREET: 225 South Lake Avenue, 9th Floor  
 CITY: Pasadena  
 STATE: CA  
 COUNTRY: USA  
 ZIP: 91101  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MB storage  
 COMPUTER: IBM compatible  
 OPERATING SYSTEM: Windows 95

SOFTWARE: Corel WordPerfect 8 version  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/863,639A  
 FILING DATE: May 28, 1997  
 CLASSIFICATION: 435  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Joseph E. Muehl  
 REGISTRATION NUMBER: 20,532  
 REFERENCE/DOCKET NUMBER: 11859-1  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (626) 796-4000  
 TELEFAX: (626) 795-6321  
 INFORMATION FOR SEQ ID NO: 94:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 24 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: Other nucleic acid  
 US-08-863-639A-94

Query Match 58.4%; Score 14.6; DB 2; Length 24;  
 Best Local Similarity 81.0%; Pred. No. 6.2e+02;  
 Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatgag 22  
 ||| ||||| ||||| ||  
 DB 2 CAGCAGCAGCAGCAGCAGCAG 22

RESULT 9

US-08-068-747-6  
 Sequence 6, Application US/08068747  
 Patent No. 5695933  
 GENERAL INFORMATION:  
 APPLICANT: Schalling, Martin  
 APPLICANT: Hudson, Thomas J.  
 APPLICANT: Housman, David E.  
 TITLE OF INVENTION: Direct Determination of Expanded  
 Nucleotide Repeats in the Human Genome  
 NUMBER OF SEQUENCES: 11  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.  
 STREET: Two Millitia Drive  
 CITY: Lexington  
 STATE: Massachusetts  
 COUNTRY: USA  
 ZIP: 02173  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: Patentin Release #1.0, Version #1.30  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/068,747  
 FILING DATE: 28-MAY-1993  
 CLASSIFICATION: 435  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Granahan, Patricia  
 REGISTRATION NUMBER: 32,227  
 REFERENCE/DOCKET NUMBER: MIT-6141  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: 617-861-6240  
 TELEFAX: 617-861-9540  
 INFORMATION FOR SEQ ID NO: 6:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 30 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: other nucleic acid  
 DESCRIPTION: /desc = "Synthetic"

US-08-068-747-6

Query Match 58.4%; Score 14.6; DB 1; Length 30;  
 Best Local Similarity 81.0%; Pred. No. 6.3e+02;  
 Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatgag 22  
 ||| ||||| ||||| ||  
 DB 1 CAGCAGCAGCAGCAGCAGCAG 21

RESULT 10

US-08-068-747-11/C  
 Sequence 11, Application US/08068747  
 Patent No. 5695933  
 GENERAL INFORMATION:  
 APPLICANT: Schalling, Martin  
 APPLICANT: Hudson, Thomas J.  
 APPLICANT: Housman, David E.  
 TITLE OF INVENTION: Direct Determination of Expanded  
 Nucleotide Repeats in the Human Genome  
 NUMBER OF SEQUENCES: 11  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.  
 STREET: Two Millitia Drive  
 CITY: Lexington  
 STATE: Massachusetts  
 COUNTRY: USA  
 ZIP: 02173  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: Patentin Release #1.0, Version #1.30  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/068,747  
 FILING DATE: 28-MAY-1993  
 CLASSIFICATION: 435  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Granahan, Patricia  
 REGISTRATION NUMBER: 32,227  
 REFERENCE/DOCKET NUMBER: MIT-6141  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: 617-861-6240  
 TELEFAX: 617-861-9540  
 INFORMATION FOR SEQ ID NO: 11:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 30 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: other nucleic acid  
 DESCRIPTION: /desc = "Synthetic"

US-08-068-747-11

Query Match 58.4%; Score 14.6; DB 1; Length 30;  
 Best Local Similarity 81.0%; Pred. No. 6.3e+02;  
 Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatgag 22  
 ||| ||||| ||||| ||  
 DB 30 CAGCAGCAGCAGCAGCAGCAG 10

RESULT 11

US-08-863-639A-30  
 Sequence 30, Application US/08863639A  
 Patent No. 5981185  
 GENERAL INFORMATION:  
 APPLICANT: Matson, Robert S.  
 APPLICANT: Coassin, Peter J.

APPLICANT: Rampal, Jang B.  
APPLICANT: Caskey, C. T.  
TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS  
NUMBER OF SEQUENCES: 95  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Sheldon & Mak  
STREET: 225 South Lake Avenue, 9th Floor  
CITY: Pasadena  
STATE: CA  
COUNTRY: USA  
ZIP: 91101  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage  
COMPUTER: IBM compatible  
OPERATING SYSTEM: Windows 95  
SOFTWARE: Corel Wordperfect 8 version  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/863,639A  
FILING DATE: May 28, 1997  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: Joseph E. Mueth  
REGISTRATION NUMBER: 20,532  
REFERENCE/DOCKET NUMBER: 11859-1  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (626) 796-4000  
TELEFAX: (626) 795-6321  
INFORMATION FOR SEQ ID NO: 30:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 30 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: Other nucleic acid  
US-08-863-639A-30

Query Match 58.4%; Score 14.6; DB 2; Length 30;  
Best Local Similarity 81.0%; Pred. No. 6.3e+02;  
Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

Oy 2 cagtagcagcacagcatgag 22  
||| ||||| ||||| ||  
Db 1 CAGCAGCAGCAGCAGCAGCAG 21

RESULT 12  
US-09-135-994-4  
Sequence 4, Application US/09135994A  
Patent No. 6280938  
GENERAL INFORMATION:  
APPLICANT: Rannum et al.  
TITLE OF INVENTION: SCAT7 GENE AND METHODS OF USE  
FILE REFERENCE: University of Minnesota  
CURRENT APPLICATION NUMBER: US/09/135,994A  
CURRENT FILING DATE: 1998-08-18  
EARLIER APPLICATION NUMBER: 60/056,170  
EARLIER FILING DATE: 1997-08-19  
NUMBER OF SEQ ID NOS: 14  
SOFTWARE: PatentIn Ver. 2.0  
SEQ ID NO 4  
LENGTH: 30  
TYPE: DNA  
ORGANISM: Homo sapiens  
US-09-135-994-4

Query Match 58.4%; Score 14.6; DB 4; Length 30;  
Best Local Similarity 81.0%; Pred. No. 6.3e+02;  
Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;  
Oy 2 cagtagcagcacagcatgag 22  
||| ||||| ||||| ||

Db 1 cagcagcagcacagcagcag 21

RESULT 13  
US-08-570-155-14  
Sequence 14, Application US/08570155  
Patent No. 5962332  
GENERAL INFORMATION:  
APPLICANT: Singer, Robert H.  
APPLICANT: Taneja, Krishan L.  
TITLE OF INVENTION: DETECTION OF TRINUCLEOTIDE REPEATS  
TITLE OF INVENTION: BY IN SITU HYBRIDIZATION  
NUMBER OF SEQUENCES: 17  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: FISH & RICHARDSON P.C.  
STREET: 225 Franklin Street  
CITY: Boston  
STATE: Massachusetts  
COUNTRY: U.S.A.  
ZIP: 02110-2804  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/570,155  
FILING DATE:  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/399,499  
FILING DATE: 07 March 1995  
PRIOR APPLICATION NUMBER: 08/214,823  
APPLICATION DATA:  
FILING DATE: 17 March 1994  
ATTORNEY/AGENT INFORMATION:  
NAME: Clark, Paul T.  
REGISTRATION NUMBER: 30,162  
REFERENCE/DOCKET NUMBER: 06353/011001  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (617) 542-5070  
TELEFAX: (617) 542-8906  
TELEX: 200154  
INFORMATION FOR SEQ ID NO: 14:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 31 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: CDNA  
US-08-570-155-14

Query Match 58.4%; Score 14.6; DB 2; Length 31;  
Best Local Similarity 81.0%; Pred. No. 6.3e+02;  
Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

Oy 2 cagtagcagcacagcatgag 22  
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Db 1 CAGCAGCAGCAGCAGCAGCAG 21

RESULT 14  
PCT-US95-02861-14  
Sequence 14, Application PC/TUS9502861  
GENERAL INFORMATION:  
APPLICANT: Singer, Robert H.  
APPLICANT: Taneja, Krishan L.  
TITLE OF INVENTION: DETECTION OF TRINUCLEOTIDE  
TITLE OF INVENTION: REPEATS  
TITLE OF INVENTION: BY IN SITU HYBRIDIZATION  
NUMBER OF SEQUENCES: 15

;; CORRESPONDENCE ADDRESS:  
;; ADDRESSEE: FISH & RICHARDSON P.C.  
;; STREET: 225 Franklin Street  
;; CITY: Boston  
;; STATE: Massachusetts  
;; COUNTRY: U.S.A.  
;; ZIP: 02110-2804  
;; COMPUTER READABLE FORM:  
;; MEDIUM TYPE: Floppy disk  
;; COMPUTER: IBM PC compatible  
;; OPERATING SYSTEM: PC-DOS/MS-DOS  
;; SOFTWARE: PatentIn Release #1.0,  
;; SOFTWARE: Version  
;; SOFTWARE: #1.30B  
;; CURRENT APPLICATION DATA:  
;; APPLICATION NUMBER: PCT/US95/02861  
;; FILING DATE: 08 March 1995  
;; CLASSIFICATION:  
;; PRIOR APPLICATION DATA:  
;; APPLICATION NUMBER: 08/214,823  
;; FILING DATE: 17 March 1994  
;; ATTORNEY/AGENT INFORMATION:  
;; NAME: Creason, Gary L.  
;; REGISTRATION NUMBER: 34,310  
;; REFERENCE/DOCKET NUMBER: 06353/010W01  
;; TELECOMMUNICATION INFORMATION:  
;; TELEPHONE: (617) 542-5070  
;; TELEFAX: (617) 542-8906  
;; TELEX: 200154  
;; INFORMATION FOR SEQ ID NO: 14:  
;; SEQUENCE CHARACTERISTICS:  
;; LENGTH: 31 base pairs  
;; TYPE: nucleic acid  
;; STRANDEDNESS: single  
;; TOPOLOGY: linear  
;; MOLECULE TYPE: CDNA  
;; PCT-US95-02861-14

Query Match 58.4%; Score 14.6; DB 5; Length 31;  
Best Local Similarity 81.0%; Pred. No. 6.3e+02;  
Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatgag 22  
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Db 1 CAGCAGCAGCAGCAGCAGCAG 21

RESULT 15  
US-08-863-639A-29/c  
; Sequence 29, Application US/08863639A  
; Patent No. 5981185  
; GENERAL INFORMATION:  
; APPLICANT: Matson, Robert S.  
; APPLICANT: Cassin, Peter J.  
; APPLICANT: Rampal, Jang B.  
; APPLICANT: Caskey, C. T.  
; TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS  
; NUMBER OF SEQUENCES: 95  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Sheldon & Max  
; STREET: 225 South Lake Avenue, 9th Floor  
; CITY: Pasadena  
; STATE: CA  
; COUNTRY: USA  
; ZIP: 91101  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage  
; COMPUTER: IBM compatible  
; OPERATING SYSTEM: Windows 95  
; SOFTWARE: Corel WordPerfect 8 version  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/863,639A

;; FILING DATE: May 28, 1997  
;; CLASSIFICATION: 435  
;; ATTORNEY/AGENT INFORMATION:  
;; NAME: Joseph E. Muehl  
;; REGISTRATION NUMBER: 20,532  
;; REFERENCE/DOCKET NUMBER: 11859-1  
;; TELECOMMUNICATION INFORMATION:  
;; TELEPHONE: (626) 796-4000  
;; TELEFAX: (626) 795-6321  
;; INFORMATION FOR SEQ ID NO: 29:  
;; SEQUENCE CHARACTERISTICS:  
;; LENGTH: 33 base pairs  
;; TYPE: nucleic acid  
;; STRANDEDNESS: single  
;; TOPOLOGY: linear  
;; MOLECULE TYPE: Other nucleic acid  
;; US-08-863-639A-29

Query Match 58.4%; Score 14.6; DB 2; Length 33;  
Best Local Similarity 81.0%; Pred. No. 6.4e+02;  
Matches 17; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 2 cagtagcagcaacagcatgag 22  
||| ||||| ||||| ||  
Db 33 CAGCAGCAGCAGCAGCAGCAG 13

Search completed: March 9, 2002, 00:54:07  
Job time: 11358 sec

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